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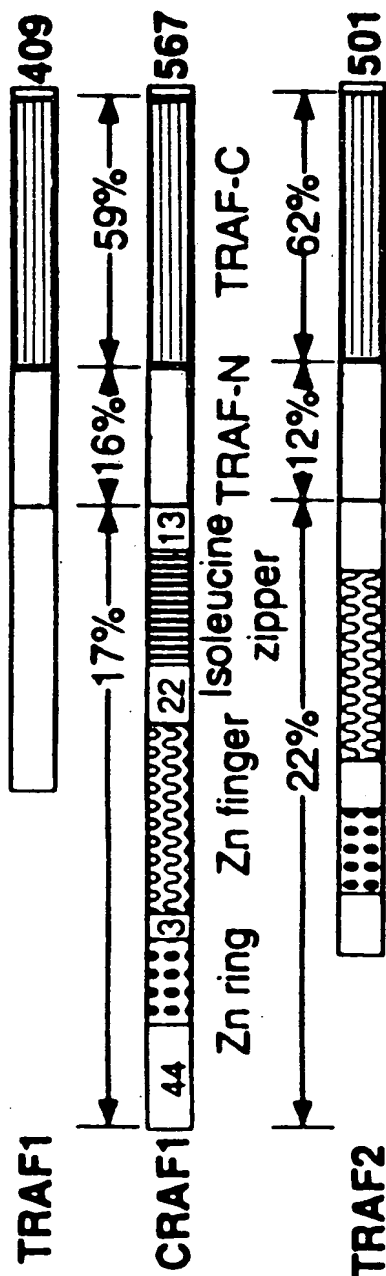
FIGURE 1

M	MESSKKMDAAGTLQPNPPLKLQPDRGAG.SVLVPEQGGYKEKFKVTVEDK	49
H	-----SP-A--T-----HT--S--TP-F-----	
M	YKCEKCRLVLCNPKQTECGHRFCESMAALLSSSSPKCTACQESI IKDKV	99
H	-----H---S-----V-----	
M	FKDNCCKREILALQVYCRNEGRGCAEQTLGHLLVHLKNECQFEELPCLR	149
H	-----I-----S-----D-H-----V-	
M	ADCCKEVLRLKDLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDCPCV	199
H	P-----A-----	
M	VVSCPHKCSVQTLRLSELSECVNAPSTCSFKRYGCVFQGTNQQIKA	249
H	-----	
M	HEASSAVQHVNLLKEWSNSLEKKVSLQNESVEKNKSIQSLHNQICSFEI	299
H	-----	
M	EIERQKEMLRNNESKILHLQRVIDSQA EKLKELDKEIRPFRQNWEEADSM	349
H	-----	
M	KSSVESLQNRVTELESVDKSAGQAARNTGLLESQLSRHDQTLVHDIRLA	399
H	-----V-----M-----	
M	DMDLRFQVLETASYNGVLIWKIRDYKRRKQEAVMGKTL SLYSQPFYTG YF	449
H	-----	
M	GYKMCARVYLNGDGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLM LMD	499
H	-----	
M	QGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQTVLENGTYIK	549
H	-----	
M	DDTIFIKVIVDTSDL PDP	567
H	-----	



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FIGURE 2A





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FIGURE 2B

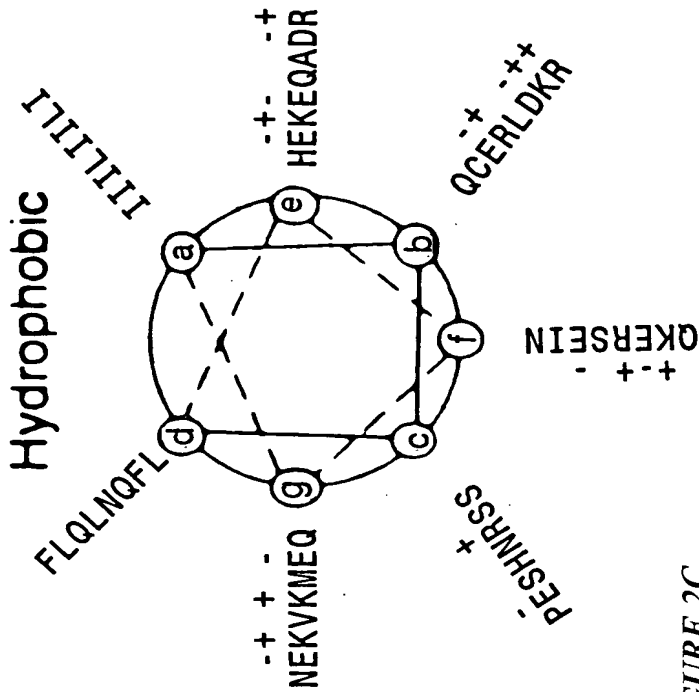


FIGURE 2D

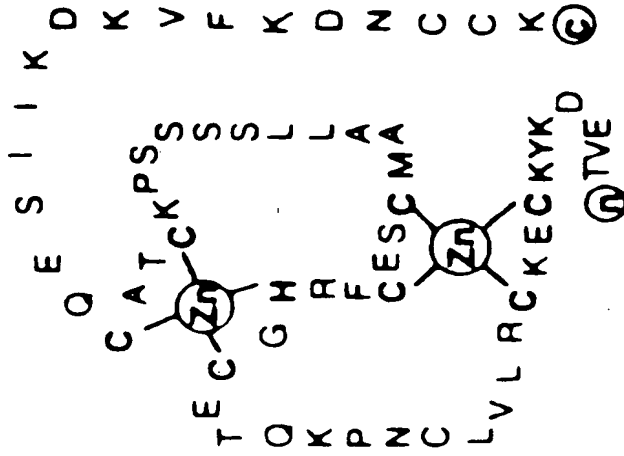
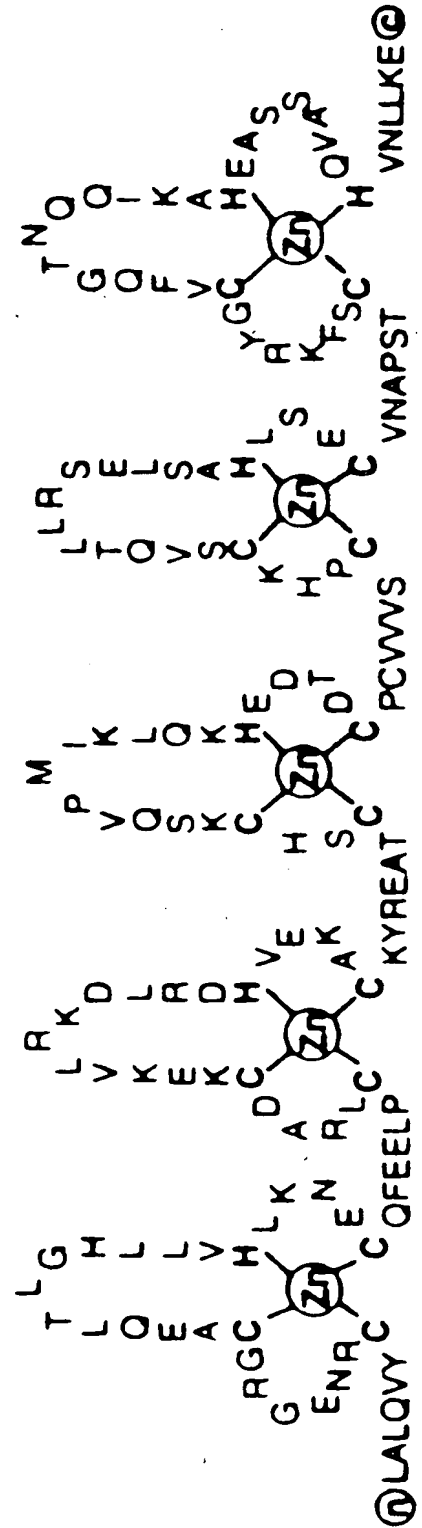
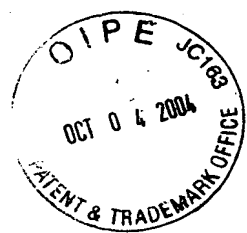







FIGURE 2C





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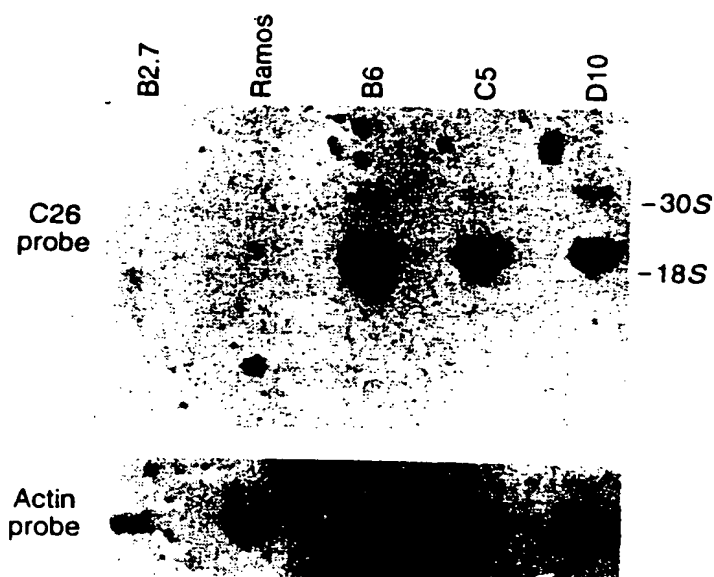
FIGURE 3

	CD40 binding	Fas binding	TNFR ^{II} binding	Homo- dimerization
 C26 (324-567)	+	-	-	+
 C26NX (324-410)	-	ND	ND	-
 C26ΔNB (324-487)	-	ND	ND	-
 TRAF domain (356-567)	+	ND	ND	+
 C26XC (415-567)	+	ND	ND	+



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FIGURE 4A





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FIGURE 4B

FIGURE 4C

FIGURE 4D

Ramos transfected with

Incubation with

pEBVHis/C26

pEBVHis/lacZ

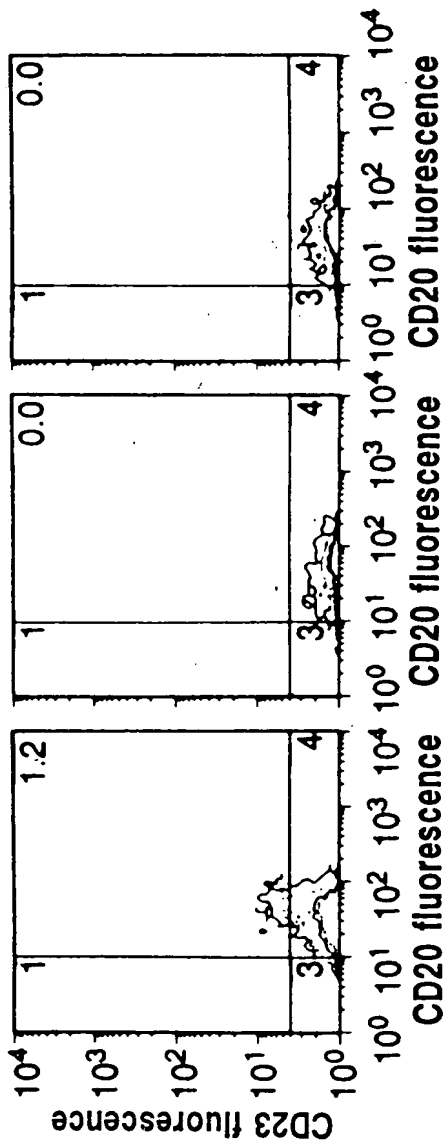
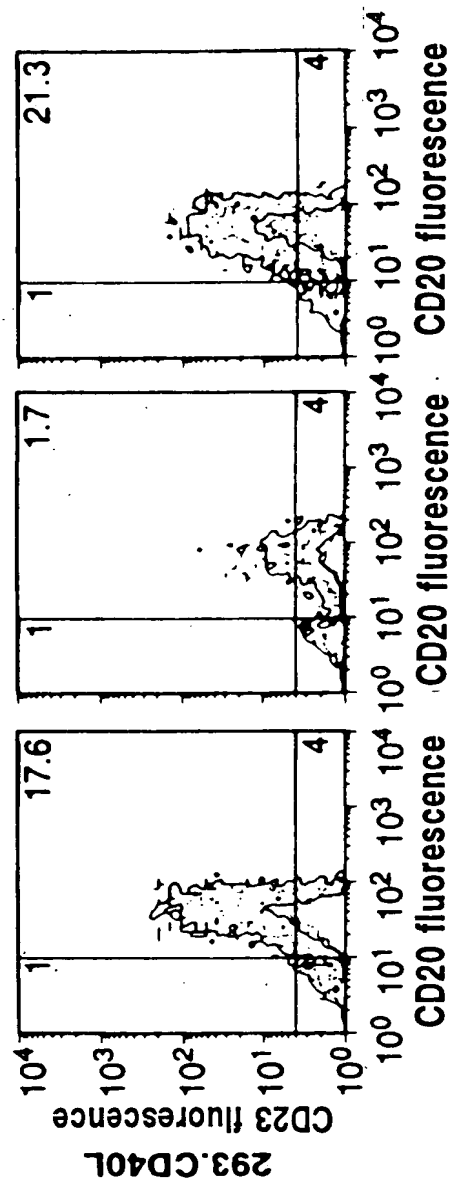


FIGURE 4E

FIGURE 4F

FIGURE 4G





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FIGURE 4H

FIGURE 4I

FIGURE 4J

Ramos transfected with

Incubation with

pEBVHis/C26

pEBVHis/lacZ

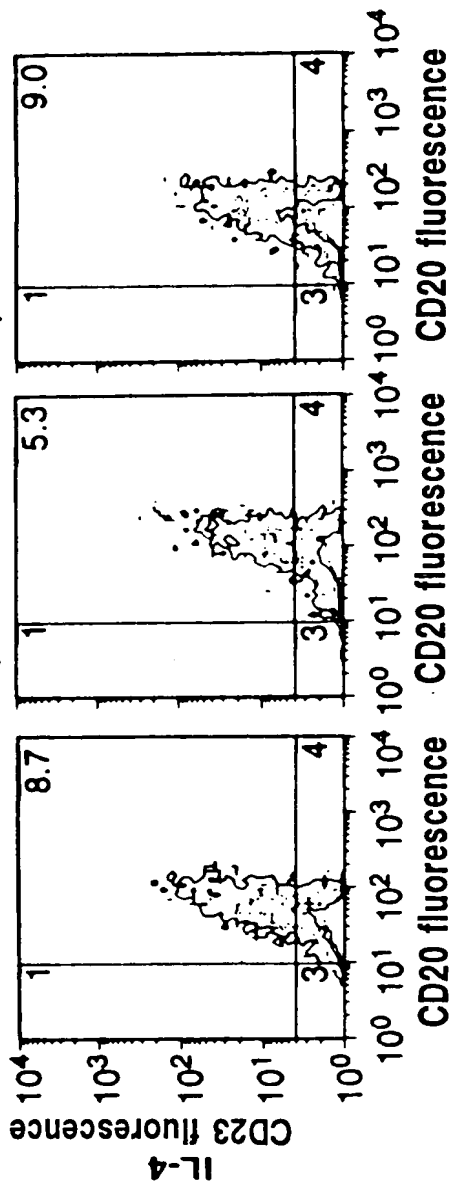
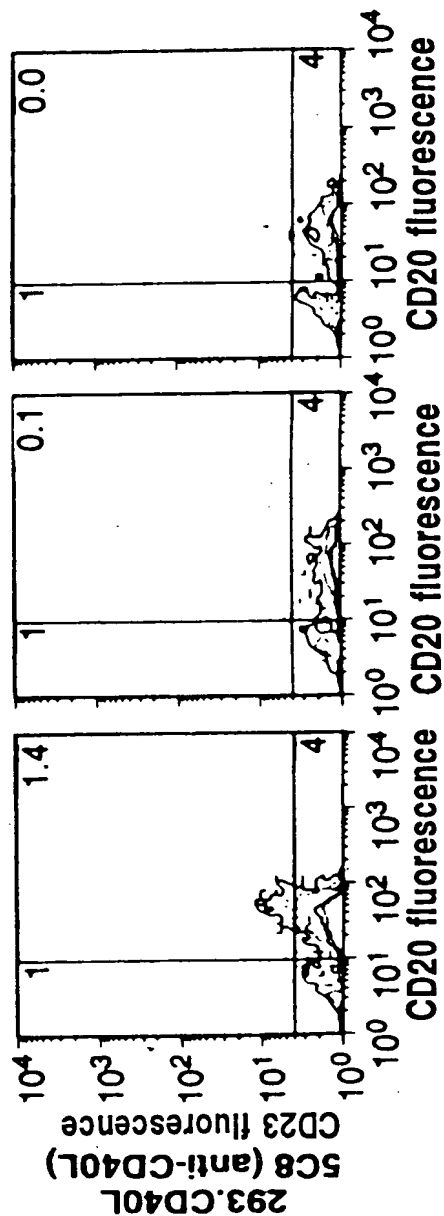


FIGURE 4K

FIGURE 4L

FIGURE 4M





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FIGURE 5A

1 GCGGCGGAG GATGCGGCG GCGCCTGAGC CGGCCGAACG GCGGGCCTCG GGGTACAGGG
61 TCCCCATTAC TTGAAGGATA AGGCTGGCAC GGCTCCGACG TCTGTGTGGA AGCTTCTCCC
121 TCCCTTCTGA GCTTCTCTAG ACTCCTTACA GCGACGGCA CAGAAATTCA GTTTCCTAAG
181 ATGGAGTCAA GCAAAAGAT GGATGCTGCT GGCACACTGC AGCCTAACCC ACCCCTAAAG
241 CTGCAGCCTG ATCGCGGCG AGGTCCTG AGGTCCTG AAGTGGCAGG CTACAAGGAG
301 AAGTTTGTGA AGACGGTGA AGACAAGTAC AAGTGGCAGG AGTGCCGCCCT GGTGCTGTGC
361 AACCCGAAGC AGACGGAGTG TGGCCACCGG TTCTGCGAGA GCTGCATGGC CGCCCTGCTG
421 AGCTCCTCCA GTCCAAAATG CACAGCGTGC CAAGAAAGCA TCATCAAAGA CAAGGTGTTT
481 AAGGATAATT GCTGCAAGAG AGAGATTCTG GCCCTTCAGG TCTACTGTG GAATGAAGGC
541 AGAGGTTGTG CGGAGCAGCT GACTCTGGGA CATCTGCTGG TGCACCTAAA AAATGAATGT
601 CAGTTTGAGG AACTTCCCTG TCTGCGTGCC GACTGCAAAG AAAAAGTACT GAGAAAGAC
661 TTGCGGGATC ACGTGGAAAA GGCCTGTAAA TACCGCGAGG CCACGTGCAG TCACTGCAAG
721 AGCCAAAGTC CCATGATCAA ACTGCAGAAA CATGAAGACA CAGATTGTCC CTGTGTGCTG
781 GTATCCTGCC CTCACAAAGT CAGCGTTCAG ACTCTTCTAA GGAGTGAGTT GAGTGCACAC
841 TTGTCCGAGT GTGTCAATGC CCCCAGCACC TGTAGTTTA AGCGCTATGG CTGCGTTT
901 CAGGGTACAA ACCAGCAGAT CAAGGCCCAT GAGGCCAGCT CCGCGGTACA GCACGTGAAC
961 CTGCTGAAGG AGTGGAGCAA CTCCCTGGAG AAGAAGGTTT CCCTGCTGCA GAATGAAGT
1021 GTTGAGAAAA ACAAGAGCAT CCAAAGCCTG CACAACCAGA TCTGCAGCTT TGAGATCGAG
1081 ATTGAGAGGC AGAAGGAGAT GCTCCGAAAC AACGAGTCCA AGATCCTTCA CCTGCAGCGG
1141 GTAATCGACA GCAAGCAGA GAAACTGAAA GAACTGGACA AGGAGATCCG TCCCTTCCGG
1201 CAGAACTGGG AGGAAGCGGA CAGCATGAAG AGCAGTGTGG AGTCCCTCCA GAACCGAGTG
1261 ACTGAGCTGG AGAGCGTAGA CAAAAGTGCG GGGCAGCGGG CTGCAACAC AGGCTTGCTG
1321 GAGTCCCAGC TGAGCCGGCA TGACCAGACG TTGAGTGTTC ATGACATCCG CTTGGCCGAC
1381 ATGGACCTGC GGTTCAGGT CCTCGAGACC GCCAGCTACA ACGGGTGCT GATCTGGAAG
1441 ATCCGTGACT ACAAGCGCGG GAAGCAGGAG GCCGTCAATGG GGAAGACCCCT GTCTCTCTAC



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FIGURE 5A

1501	AGCCAGCCTT	TCTACACAGG	TTATTTTGGC	TATAAGATGT	GTGCCAGGGT	CTACCTGAAT
1561	GGGACGGAA	TGGGAAAGG	GACACACTTG	TCGCTGTTT	TTGTCATTAT	GCGTGGAGAA
1621	TATGATGCTC	TGTTGCCATG	GCCGTTCAAG	CAGAAAGTGA	CACTTATGCT	GATGGATCAG
1681	GGGTCCTCTC	GCCGTCATCT	GGGAGATGCG	TTCAAGCCTG	ACCCCAACAG	CAGCAGCTTC
1741	AAGAAACCCA	CCGGAGAGAT	GAATATCGCC	TCTGGCTGCC	CAGTCTTTGT	CGCCCCAACT
1801	GTTCTAGAGA	ACGGGACGTA	TATTAAGAT	GATACAATCT	TTATTAAGGT	CATAGTGGAT
1861	ACCTCGGATC	TGCCGTACCC	CTGACAAGAA	AGCAGGGCGG	TGGATTCAGC	AGAAGGTAAC
1921	TCCTCTGGGG	GGGTGAGCTA	GTGCTTTCAC	GGAGGTCCTC	GCCCTCAGAA	AGGACCTTGT
1981	GGCGCAGAGG	AAGCAGCCCG	AGGAGGAGAA	GGAGGTCGAG	TGGCTGGCAG	GAGAGCCACA
2041	TGTGAAAACA	GACCCCAACG	GATTTTCTAA	TAAACTAGCC	ACACCCACTC	TGAAGGATTA
2101	TTTATCCATC	AACAAGATAA	ATACTGCTGT	CAGAGAAGGT	TTTCATTTTC	ATTTTAAAG
2161	ATCTAGTATT	AAGGTGGGAA	CATATATGCT	AAAAGAAAC	ATGATTTTTC	TTCCCTTAAC
2221	TAAACACCAA	AAAGAGAACA	CATGTGGGG	TAGCTGGAGT	GTGTACAGTA	CCTCGAGGGC
2281	TTAAAATCAT	AAACAATCAC	ATACTCATCC	TAAAATTTCAG	GGTGCAACTC	CGTTTCAAAT
2341	ATTGTATATT	GTCTATTTA				



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FIGURE 6A

1 CGGGGGAGCG CGGCGGGGCC GCGCGGTGCG CGAGCCGGGG TTGCAGCCCA GCCGGGACTT
61 TCCAGCCGGC GGCAGCCGCG GCGGTCGTGG GCTCTTCCCC GCCCCCGTC ATGGGGCAGC
121 CCGGGAGCA GAACGCTGCG GACCGGGCGG GAGGACGCG CCGGCGCCC TGAGCCGGCC
181 GAGCGGCAC GGACCGCGAG AACTCCTCTT TCCTAAATG GAGTCAGTA AAAAGATGGA
241 CTCCTCTGGC GCGCTGCAGA CTAACCCGCC GCTAAAGCTG CACACTGACC GTAGTGCTGG
301 GACGCCAGTT TTTGTCCCTG AACAGGAGG TTACAAGGAA AAGTTGTGA AGACCGTGGA
361 GGACAAAGTAC AAGTGTGAGA AGTGCCACCT GGTGCTGTGC AGCCCGAAGC AGACCGAGTG
421 TGGGCACCGC TTCTGCGAGA GCTGCATGGC GGGCCTGCTG AGCTCTTCAA GTCCAAAATG
481 TACAGCGTGT CAAGAGAGCA TCGTTAAAGA TAAGGTGTTT AAGGATAATT GCTGCAAGAG
541 AGAAATTCTG GCTCTTCAGA TCTATTGTCG GAATGAAAGC AGAGGTTGTG CAGAGCAGTT
601 AACGCTGGG CATCTGCTGG TGCATTTAAA AAATGATTGC CATTTTGAAG AACTTCCATG
661 TGTGCGTCCT GACTGCAAAG AAAAGGTCTT GAGGAAAGAC CTGCGAGACC ACGTGGAGAA
721 GCGGTGTAAT TACCGGAAG CCACATGCAG CCACTGCAAG AGTCAGGTTT CGATGATCGC
781 GCTGCAGAAA CACGAAGACA CCGACTGTCC CTGCGTGGTG GTGTCTGCTG CACACAAGTG
841 CAGCGTCCAG ACTCTCCTGA GGAGCGAGTT GAGTGCACAC TTGTCAGAGT GTGTCAATGC
901 CCCAGCACC TGTAGTTTAA AGCGTATGG CTGCGTTTTT CAGGGACAA ACCAGCAGAT
961 CAAGGCCAC GAGGCCAGCT CCGCCGTGCA GCACGTCAAC CTGCTGAAGG AGTGGAGCAA
1021 CTCGCTCGAA AAGAAAGTTT CCTTGTGCA GAATGAAAGT GTAGAAAAA ACAAGAGCAT
1081 ACAAGTTTG CACAATCAGA TATGTAGCTT TGAAATTGAA ATTGAGAGAC AAAAGGAAAT
1141 GCTTCGAAAT AATGAATCCA AAATCCTTCA TTTACAGCGA GTGATCGACA GCCAAGCAGA
1201 GAAACTGAAG GAGCTTGACA AGGAGATCCG GCGCTTCCGG CAGAACTGGG AGGAAGCAGA
1261 CAGCATGAAG AGCAGCGTGG AGTCCCTCCA GAACCGCGTG ACCGAGCTGG AGAGCGTGG
1321 CAAGAGTGCG GGGCAAGTGG CTCGGAAACAC AGGCCTGCTG GAGTCCCAGC TGAGCCGGCA
1381 TGACCAGATG CTGAGTGTGC ACGACATCCG CCTAGCCGAC ATGGACCTGC GCTTCCAGGT
1441 CCTGGAGACC GCCAGCTACA ATGGAGTGCT CATCTGGAAG ATTCCGGACT ACAAGCGGCG



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FIGURE 6B

1501	GAAGCAGGAG	GCCGTCATGG	GGAAGACCCCT	GTCCCTTTAC	AGCCAGCCCTT	TCTACACTGG
1561	TTACTTTGGT	TATAAGATGT	GTGCCAGGGT	CTACCTGAAC	GGGACGGGA	TGGGGAAGGG
1621	GACGCACTTG	TCGCTGTTTT	TTGTCATCAT	GCGTGGAGAA	TATGATGCCC	TGCTTCCCTTG
1681	GCCGTTTAAG	CAGAAAGTGA	CACATCATGCT	GATGGATCAG	GGGTCCTCTC	GACGTCAATT
1741	GGGAGATGCA	TTCAAGCCCG	ACCCCAACAG	CAGCAGCTTC	AGAAAGCCCA	CTGGAGAGAT
1801	GAATATCGCC	TCTGGCTGCC	CAGTCTTTGT	GGCCCAAACT	GTTCTAGAAA	ATGGGACATA
1861	TATTAAGAT	GATACAAATT	TTATTAAAGT	CATAGTGGAT	ACTTCGGATC	TGCCCCGATCC
1921	CTGATAAGTA	GCTGGGGAGG	TGGATTTAGC	AGAAGGCAAC	TCCTCTGGGG	GATTTGAACC
1981	GGTCTGTCTT	CAC TGAGGTC	CTCGCGCTCA	GAAAAGGACC	TTGTGAGACG	GAGGAAGCGG
2041	CAGAAGGCGG	ACGCGTGCCG	GCGGGAGGAG	CCACGCGTGA	GCACACCTGA	CACGTTTAT
2101	AATAGACTAG	CCACACTTCA	CTCTGAAGAA	TTATTTATCC	TTCAACAAGA	TAAATATTGC
2161	TGTCAGAGAA	GGTTTTTCATT	TTCATTTTAA	AAGATCTAGT	TAATTAAGGT	GGAAAACATA
2221	TATGCTAAAC	AAAAGAAACA	TGATTTTCT	TCCTTAAACT	TGAACACCAA	AAAAACACAC
2281	ACACACACAC	ACGTGGGGAT	AGCTGGACAT	GTCAGCATGT	TAAGTAAAAG	GAGAAATTTAT
2341	GAATAAGTAA	TGCAATTCTG	ATATCTTCTT	TCTAAAATTC	AAGAGTGCAA	TTTGTGTTTCA
2401	AATACAGTAT	ATTGTCTATT	TTTAAGGCCT	CCAAAAAAA	AAAAAATTC	GGCCG